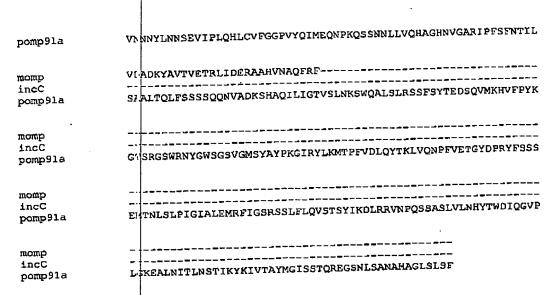
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! Sequence:	i a a C	
! Sequence:	incc	
! Sequence:	bombaj	<u>.</u> .
CLUSTAL W ()	. 831 a	ultiple sequence alignment
CHOSIND W /-		F •
momp		
incC		
	MI	QMRLWGFLFLSSFCQV5YLRANDVLLPLSGIH3GEDLELFTLR59SPTKTTYSLRKDF
pomp91a	PII	principal de la company de la
		www.i.kgvi.vear.ssassi.oalpvg
momp	<b></b>	
		MTSPIPFQ-
incC	TI	CDFAGNSIHKPGAAFLNLKGDLFFINSTPLAALTFKNIHLGARGAGLFSESNVTFKGL
pomp91a	1.	CDEAGNOTING COLUMN COLU
		NPAEPSLMIDGILWEGFGGDPCDPCATWCDAISMRVGYYGDFVF
momp	_	NPAEPSIMIDGILWEGEGGGECOTT THANKITONI SETTI OO
	_	
incC		LVLENNESWGGVLTTSGDLSFINNTSVLCONNISYGPGGALLLQGRKSKALFFRDNRG
pomp9la	H.	1: *:: : : : : : : : : : : : : : : :
#4WD	D	RVLKTDVNKEFQMGAKPTTDTGNSAAPS
momp		
incC	~	RDRLPTASIILQVGGAPTGGAGAPTQFG ILFLKNKAVNQDESHPGYGGAVSSISPGSPITFADNQEILFQENEGELGGAIYNDQGAI
pomp9la	T	ILFHANKAVNODESHI GIGGOVE
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		FDVFCTLGATSGYLKGNSASFNLVGLFGDNENQKTVKAESVFNMSFDQSVVELYTDTT
	-	-FDVFCTLGATSGYLKGNSASFNLVGLEGDNENGATVALGSVETGVLFQVALLMQGETN -STLQQSTKGARTGVLVVTAILMTISLLAIIIILAVLGFTGVLFQVALLMQGETN -STLQQSTKGARTGVLVVTAILMTISLEAN FINNSALGLNGGATYMQATGSILRLHAN
momp	_	LSTIOOSTKGARTGVLVVTAILMTISLLAIIIIILAVLGFTG-TOOGATGCILBIHAN
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pomp9la	1.	G314EBR021200
momp	-	T TWAMVSGSTTCFIALIG
_		
incC		GDIEFCGNKVRSQFHSHINSTSNFTNNAITIQGAPREFSLSANEGHRICFYDPI1SATE
pomp91a	`	
momp		
incC		INOPVRICEGVISIEGGAIL
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DOMD 2 + F		
		- ASFQYAQSKPKVEELN VICTORIAL
qmom	•	TLGLILTNKNTPLPAS
incC		TECCHISEROKI VITNLGFNLENLDSSDPARIRATEKASI
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Doutharn		
•••		TINKPKGYVGKEFPLDLTAGTDAATG
momp		FINAPAGIVONE
incC		STEEN SKRYTTSTILSAKKLYTAPSRPEKDIQNLI LAESEIM
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501-5-2-		
		TKDASIDYHEWQASLALSYRLNMFTPYIGVKWSRASFDADTIRIA
		TKDASIDYHEWQASLALSYKLNMFTF1IGVANGTEN
quon		TRUASIDI NOMENO DE LA CONTRACTO DE LA CONTRACTO DE LA CONTRACTOR DE LA CON
incC	٠	YGYQGSWEFSWSPNDTKEXKTIIASWTPTGEFSLDPKRRGSFIPTTLWSTFSGLNIASN
pomp9la		ACAOCEMER 2M2 RULL VETT 1 TO 1
h		
		QPKSATAIFDTTTLNPTIAGAGDVKTGAEGQLGDTMQIVSLQLNKMKSRKSCGIAVGTT
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momp		Zr vo
incC		



<sup>&</sup>quot;\*" means that the residues or nucleotides in that column are identical in all sequences in the alignment

<sup>&</sup>quot;:" means that conserved substitutions have been observed

<sup>&</sup>quot;" means that senti-conserved substitutions are observed

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947 aa
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Murdin,A.D., Dunn,P.L. and Oomen,R.P.
Nucleic acid molecules encoding POMP91A protein of Chlamydia
Patent: US 6693087-A 3 17-FEB-2004;
Aventis Pasteur Limited; Toronto;
REFERENCE
     AUTHORS
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      JOURNAL
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Murdin,A.O., Dunn,P.L. and Oomen,R.P.

Nucleic adid molecules encoding inclusion membrane protein C of
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stephens momp
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                                                                                                            linear
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Created: May 30, 2000.

sequence updated: Nov 1, 1996.

annotation updated: May 2, 2006.

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AAC68276.1, H71484

xrefs (non-sequence databases): PHCI-2DPAGE:Q46409,

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                       Sayada,C. Denamur,E. and Elion,J. Complete sequence of the major outer membrane protein-encoding gene of chlamydia trachomatis serovar Da Gene 120 (1), 129-130 (1992)
REFERENCE
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                        Stothard, D.R., Boguslawski, G. and Jones, R.B.
Stothard, D.R., Boguslawski, G. and Jones, R.B.
Phylogenetic analysis of the Chlamydia trachomatis major outer membrane protein and examination of potential pathogenic
 REFERENCE
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9673241
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 REFERENCE
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                         Koonin,E.V. and Davis,R.W.
Genome sequence of an obligate intracellular pathogen of humans:
Chlamydia trachomatis
Science 282 (5389), 754-759 (1998)
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NUCLEOTIDE SEQUENCE [LARGE SCALE GENOMIC DNA].
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ON SEP 27, 2005 this sequence version replaced gi:7442973.
[FUNCTION] Structural rigidity of the outer membrane of elementary bodies and porin forming, permitting diffusion of solutes through the intracellular reticulate body membrane.
[SUBUNIT] Disulfide bond interactions within and between MOMP molecules and other components form high molecular-weight oligomers.
     JOURNAL
       PUBMED
      REMARK
  COMMENT
                         oligomers.
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